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# Exploratory data analysis (EDA) using python: a tutorial

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# EXPLORATORY DATA ANALYSIS (EDA) USING PYTHON: A TUTORIAL

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OCT., 2024

# AGENDA

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- **Introduction**
- **Overview**
- **Data Inspection & Cleaning Steps**
- **Graphical EDA**
- **Conclusions**
- **References**

# INTRODUCTION: WHAT IS EDA?

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Exploratory data analysis is an analysis technique to analyze and investigate the data set and summarize the main characteristics of the dataset. Main advantage of EDA is providing the data visualization of data after conducting the analysis.

Tukey defined data analysis in 1961 as: "Procedures for analyzing data, techniques for interpreting the results of such procedures, ways of planning the gathering of data to make its analysis easier, more precise or more accurate, and all the machinery and results of (mathematical) statistics which apply to analyzing data".

[https://en.wikipedia.org/wiki/Exploratory\\_data\\_analysis](https://en.wikipedia.org/wiki/Exploratory_data_analysis)

# OVERVIEW

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Obesity is a complex disease involving having too much body fat. Obesity isn't just a cosmetic concern. It's a medical problem that increases the risk of many other diseases and health problems. These can include heart disease, diabetes, high blood pressure, high cholesterol, liver disease, sleep apnea and certain cancers.

There are many reasons why some people have trouble losing weight. Often, obesity results from inherited, physiological and environmental factors, combined with diet, physical activity and exercise choices.

The good news is that even modest weight loss can improve or prevent the health problems associated with obesity. A healthier diet, increased physical activity and behavior changes can help you lose weight. Prescription medicines and weight-loss procedures are other options for treating obesity. In this Exploratory Data Analysis (EDA) using Python we will examine the NObeyesdad (Obesity level of the individual) as a function of several factors (i.e., age, gender, height, alcohol consumption ..etc.).

<https://www.mayoclinic.org/diseases-conditions/obesity/symptoms-causes/syc-20375742>

# DATASET INFORMATION

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This dataset include data for the estimation of obesity levels in individuals from the countries of Mexico, Peru and Colombia, based on their eating habits and physical condition. The data contains 17 attributes and 2111 records, the records are labeled with the class variable NObesity (Obesity Level), that allows classification of the data using the values of Insufficient Weight, Normal Weight, Overweight Level I, Overweight Level II, Obesity Type I, Obesity Type II and Obesity Type III. 77% of the data was generated synthetically using the Weka tool and the SMOTE filter, 23% of the data was collected directly from users through a web platform.

<https://archive.ics.uci.edu/dataset/544/estimation+of+obesity+levels+based+on+eating+habits+and+physical+condition>

# DATA INSPECTION & CLEANING STEPS

---

1. Dataset dimensions
2. Titles of columns
3. Data types
4. Missing values
5. Nulls in the dataset (not required for this dataset)
6. Duplicate rows
7. NaN values
8. Infinity values
9. Outliers detection
10. Encode categorical features



# LOADING PYTHON LIBRARIES

```
import matplotlib.pyplot as plt
import seaborn as sns
import warnings
import pandas as pd
import numpy as np
from sklearn.preprocessing import OneHotEncoder, StandardScaler, LabelEncoder
from sklearn.model_selection import train_test_split
from sklearn.metrics import confusion_matrix
warnings.filterwarnings("ignore")
```

**pandas (<https://pandas.pydata.org>)**

**numpy (<https://numpy.org>)**

**matplotlib (<https://matplotlib.org>)**

**seaborn (<https://seaborn.pydata.org>)**

**sklearn (<https://scikit-learn.org/stable>)**



# DATA SET ATTRIBUTES

1. Age: The age of the individual.
  2. Gender: The gender of the individual (e.g., Male, Female).
  3. Height: The height of the individual in meters.
  4. Weight: The weight of the individual in kilograms.
  5. CALC: Unknown column. You might need to check the data source or documentation to understand what this column represents.
  6. FAVC: Whether the individual frequently consumes high caloric food (e.g., yes, no).
  7. FCVC: Frequency of consumption of vegetables (numeric scale).
  8. NCP: Number of main meals per day (numeric scale).
  9. SCC: Squamous cell carcinoma
  10. SMOKE: Whether the individual smokes (e.g., yes, no).
  11. CH2O: Consumption of water daily (numeric scale).
  12. family\_history\_with\_overweight: Whether the individual has a family history of overweight (e.g., yes, no).
  13. FAF: Physical activity frequency (numeric scale).
  14. TUE: Time using technology devices (numeric scale).
  15. CAEC: Unknown column. You might need to check the data source or documentation to understand what this column represents.
  16. MTRANS: Mode of transportation (e.g., Public Transportation, Walking).
  17. NObesidad: Obesity level of the individual (e.g., Normal\_Weight, Overweight\_Level\_I).
- Note:** The data set contains a mixture of numeric and categorical data.

# PANDAS FUNCTIONS FOR DATA INSPECTION

---

- `df.head()/df.tail()`
- `df.sample()`
- `df.info()`
- `df.columns`
- `df.describe()`
- `df.shape`
- `Df.count()`

<https://www.geeksforgeeks.org/pandas-functions-in-python/>

# USING PANDAS FOR LOADING THE DATA FILE & VIEWING THE FIRST 5 ROWS

```
data = pd.read_csv("/content/sample_data/ObesityDataSet_raw_and_data.csv") # Load data set using pandas
data.head(5)
```

	Age	Gender	Height	Weight	CALC	FAVC	FCVC	NCP	SCC	SMOKE	CH2O	family_history_with_overweight	FAF	TUE	(
0	21.0	Female	1.62	64.0	no	no	2.0	3.0	no	no	2.0	yes	0.0	1.0	Some
1	21.0	Female	1.52	56.0	Sometimes	no	3.0	3.0	yes	yes	3.0	yes	3.0	0.0	Some
2	23.0	Male	1.80	77.0	Frequently	no	2.0	3.0	no	no	2.0	yes	2.0	1.0	Some
3	27.0	Male	1.80	87.0	Frequently	no	3.0	3.0	no	no	2.0	no	2.0	0.0	Some
4	22.0	Male	1.78	89.8	Sometimes	no	2.0	1.0	no	no	2.0	no	0.0	0.0	Some

**Dataset source: UC Irvine-Machine Learning Repository**

<https://archive.ics.uci.edu/dataset/544/estimation+of+obesity+levels+based+on+eating+habits+and+physical+condition>

# EXAMINING 5 ROWS IN RANDOM & AT THE END OF THE DATASET

1 data.sample(5)

	Gender	Age	Height	Weight	family_history_with_overweight	FAVC	FCVC	NCP	CAEC	SMOKE	CH2O	SCC	FAF	
1679	Male	31.761799	1.751688	119.205308	yes	yes	2.149610	3.000000	Sometimes	no	2.133876	no	0.393452	0.3
752	Male	21.142432	1.855353	86.413388	yes	yes	2.000000	3.000000	Sometimes	no	1.345298	no	1.097905	1.0
1546	Male	25.298400	1.827279	120.996074	yes	yes	3.000000	3.000000	Sometimes	no	3.000000	no	1.110215	0.3
1741	Male	28.255199	1.816547	120.699119	yes	yes	2.997951	3.000000	Sometimes	no	2.715856	no	0.739881	0.3
1790	Male	23.147644	1.815514	120.337664	yes	yes	2.996717	2.791366	Sometimes	no	2.626309	no	1.194898	0.0

1 data.tail(5)

	Gender	Age	Height	Weight	family_history_with_overweight	FAVC	FCVC	NCP	CAEC	SMOKE	CH2O	SCC	FAF	TUE
2106	Female	20.976842	1.710730	131.408528	yes	yes	3.0	3.0	Sometimes	no	1.728139	no	1.676269	0.906247
2107	Female	21.982942	1.748584	133.742943	yes	yes	3.0	3.0	Sometimes	no	2.005130	no	1.341390	0.599270
2108	Female	22.524036	1.752206	133.689352	yes	yes	3.0	3.0	Sometimes	no	2.054193	no	1.414209	0.646288
2109	Female	24.361936	1.739450	133.346641	yes	yes	3.0	3.0	Sometimes	no	2.852339	no	1.139107	0.586035
2110	Female	23.664709	1.738836	133.472641	yes	yes	3.0	3.0	Sometimes	no	2.863513	no	1.026452	0.714137

# USING PANDA FUNCTIONS FOR VIEWING THE DATA

```
In [136]: data.columns
```

```
Out[136]: Index(['Age', 'Gender', 'Height', 'Weight', 'CALC', 'FAVC', 'FCVC', 'NCP',  
                'SCC', 'SMOKE', 'CH20', 'family_history_with_overweight', 'FAF', 'TUE',  
                'CAEC', 'MTRANS', 'NObeyesdad'],  
              dtype='object')
```

```
In [138]: data.shape # Fetch the data set dimensions before cleaning
```

```
Out[138]: (2111, 17)
```

**Data set size:**  
**2111 rows x 17 columns**

```
data.info() # check data type in each column
```

```
<class 'pandas.core.frame.DataFrame'>
```

```
RangeIndex: 2111 entries, 0 to 2110
```

```
Data columns (total 17 columns):
```

#	Column	Non-Null Count	Dtype
0	Age	2111 non-null	float64
1	Gender	2111 non-null	object
2	Height	2111 non-null	float64
3	Weight	2111 non-null	float64
4	CALC	2111 non-null	object
5	FAVC	2111 non-null	object
6	FCVC	2111 non-null	float64
7	NCP	2111 non-null	float64
8	SCC	2111 non-null	object
9	SMOKE	2111 non-null	object
10	CH20	2111 non-null	float64
11	family_history_with_overweight	2111 non-null	object
12	FAF	2111 non-null	float64
13	TUE	2111 non-null	float64
14	CAEC	2111 non-null	object
15	MTRANS	2111 non-null	object
16	NObeyesdad	2111 non-null	object

```
dtypes: float64(8), object(9)
```

```
memory usage: 280.5+ KB
```



# DATA STATISTICS & MISSING VALUES CHECK

```
data.describe()
```

	Age	Height	Weight	FCVC	NCP	CH2O	FAF	TUE
count	2111.000000	2111.000000	2111.000000	2111.000000	2111.000000	2111.000000	2111.000000	2111.000000
mean	24.312600	1.701677	86.586058	2.419043	2.685628	2.008011	1.010298	0.657866
std	6.345968	0.093305	26.191172	0.533927	0.778039	0.612953	0.850592	0.608927
min	14.000000	1.450000	39.000000	1.000000	1.000000	1.000000	0.000000	0.000000
25%	19.947192	1.630000	65.473343	2.000000	2.658738	1.584812	0.124505	0.000000
50%	22.777890	1.700499	83.000000	2.385502	3.000000	2.000000	1.000000	0.625350
75%	26.000000	1.768464	107.430682	3.000000	3.000000	2.477420	1.666678	1.000000
max	61.000000	1.980000	173.000000	3.000000	4.000000	3.000000	3.000000	2.000000

## Dataset Statistics Summary

```
# Check for missing values in the DataFrame
missing_values = data.isnull().sum()
print("Missing Values:")
print(missing_values)
```

```
Missing Values:
Age                                0
Gender                            0
Height                            0
Weight                            0
CALC                              0
FAVC                              0
FCVC                              0
NCP                               0
SCC                               0
SMOKE                             0
CH2O                              0
family_history_with_overweight    0
FAF                                0
TUE                               0
CAEC                              0
MTRANS                            0
NObeyesdad                        0
dtype: int64
```

As seen, there are no missing values in this dataset

## Missing Values Check

## No missing values

# INSPECTING THE DATASET FOR DUPLICATE ROWS & DROPPING THE DUPLICATE ROWS

```
# Check for duplicate rows in the data set
duplicate_rows = data.duplicated().sum()
print("Duplicate Rows:")
print(duplicate_rows)
```

Duplicate Rows:  
24

As shown above, there is 24 duplicate rows in the data set.

**24 duplicate row**  
**After dropping the 24**  
**duplicate Rows, new dataset**  
**size:**  
**2087 rows x 17 columns**

```
data=data.drop_duplicates() # use dataframe.drop_duplicates() to drop the duplicate rows
display(data)
```

	Age	Gender	Height	Weight	CALC	FAVC	FCVC	NCP	SCC	SMOKE	CH2O	family_history_with_overweight
0	21.000000	Female	1.620000	64.000000	no	no	2.0	3.0	no	no	2.000000	yes
1	21.000000	Female	1.520000	56.000000	Sometimes	no	3.0	3.0	yes	yes	3.000000	yes
2	23.000000	Male	1.800000	77.000000	Frequently	no	2.0	3.0	no	no	2.000000	yes
3	27.000000	Male	1.800000	87.000000	Frequently	no	3.0	3.0	no	no	2.000000	no
4	22.000000	Male	1.780000	89.800000	Sometimes	no	2.0	1.0	no	no	2.000000	no
...	...	...	...	...	...	...	...	...	...	...	...	...
2106	20.976842	Female	1.710730	131.408528	Sometimes	yes	3.0	3.0	no	no	1.728139	yes
2107	21.982942	Female	1.748584	133.742943	Sometimes	yes	3.0	3.0	no	no	2.005130	yes
2108	22.524036	Female	1.752206	133.689352	Sometimes	yes	3.0	3.0	no	no	2.054193	yes
2109	24.361936	Female	1.739450	133.346641	Sometimes	yes	3.0	3.0	no	no	2.852339	yes
2110	23.664709	Female	1.738836	133.472641	Sometimes	yes	3.0	3.0	no	no	2.863513	yes

2087 rows × 17 columns

```
data.shape # check data set dimension after removing duplicates
```

(2087, 17)



# NAN VALUES CHECK

```
# Check if the set has NaN values
print(data.isna())
```

	Age	Gender	Height	Weight	CALC	FAVC	FCVC	NCP	SCC	SMOKE
0	False	False	False	False	False	False	False	False	False	False
1	False	False	False	False	False	False	False	False	False	False
2	False	False	False	False	False	False	False	False	False	False
3	False	False	False	False	False	False	False	False	False	False
4	False	False	False	False	False	False	False	False	False	False
...	...	...	...	...	...	...	...	...	...	...
2106	False	False	False	False	False	False	False	False	False	False
2107	False	False	False	False	False	False	False	False	False	False
2108	False	False	False	False	False	False	False	False	False	False
2109	False	False	False	False	False	False	False	False	False	False
2110	False	False	False	False	False	False	False	False	False	False

	CH2O	family_history_with_overweight	FAF	TUE	CAEC	MTRANS
0	False	False	False	False	False	False
1	False	False	False	False	False	False
2	False	False	False	False	False	False
3	False	False	False	False	False	False
4	False	False	False	False	False	False
...	...	...	...	...	...	...
2106	False	False	False	False	False	False
2107	False	False	False	False	False	False
2108	False	False	False	False	False	False
2109	False	False	False	False	False	False
2110	False	False	False	False	False	False

	NObeyesdad
0	False
1	False
2	False
3	False
4	False
...	...
2106	False
2107	False
2108	False
2109	False
2110	False

[2087 rows x 17 columns]

No NaN values in the dataset

# INFINITY VALUES CHECK

```
# Check the presence of infinity values in the data set
data.isin([np.inf, -np.inf])
print(data)
```

No  $\pm\infty$  values in  
the data set

	Age	Gender	Height	Weight	CALC	FAVC	FCVC	NCP	\
0	21.000000	Female	1.620000	64.000000	no	no	2.0	3.0	
1	21.000000	Female	1.520000	56.000000	Sometimes	no	3.0	3.0	
2	23.000000	Male	1.800000	77.000000	Frequently	no	2.0	3.0	
3	27.000000	Male	1.800000	87.000000	Frequently	no	3.0	3.0	
4	22.000000	Male	1.780000	89.800000	Sometimes	no	2.0	1.0	
...	...	...	...	...	...	...	...	...	
2106	20.976842	Female	1.710730	131.408528	Sometimes	yes	3.0	3.0	
2107	21.982942	Female	1.748584	133.742943	Sometimes	yes	3.0	3.0	
2108	22.524036	Female	1.752206	133.689352	Sometimes	yes	3.0	3.0	
2109	24.361936	Female	1.739450	133.346641	Sometimes	yes	3.0	3.0	
2110	23.664709	Female	1.738836	133.472641	Sometimes	yes	3.0	3.0	

	SCC	SMOKE	CH20	family_history_with_overweight	FAF	TUE	\
0	no	no	2.000000	yes	0.000000	1.000000	
1	yes	yes	3.000000	yes	3.000000	0.000000	
2	no	no	2.000000	yes	2.000000	1.000000	
3	no	no	2.000000	no	2.000000	0.000000	
4	no	no	2.000000	no	0.000000	0.000000	
...	...	...	...	...	...	...	
2106	no	no	1.728139	yes	1.676269	0.906247	
2107	no	no	2.005130	yes	1.341390	0.599270	
2108	no	no	2.054193	yes	1.414209	0.646288	
2109	no	no	2.852339	yes	1.139107	0.586035	
2110	no	no	2.863513	yes	1.026452	0.714137	

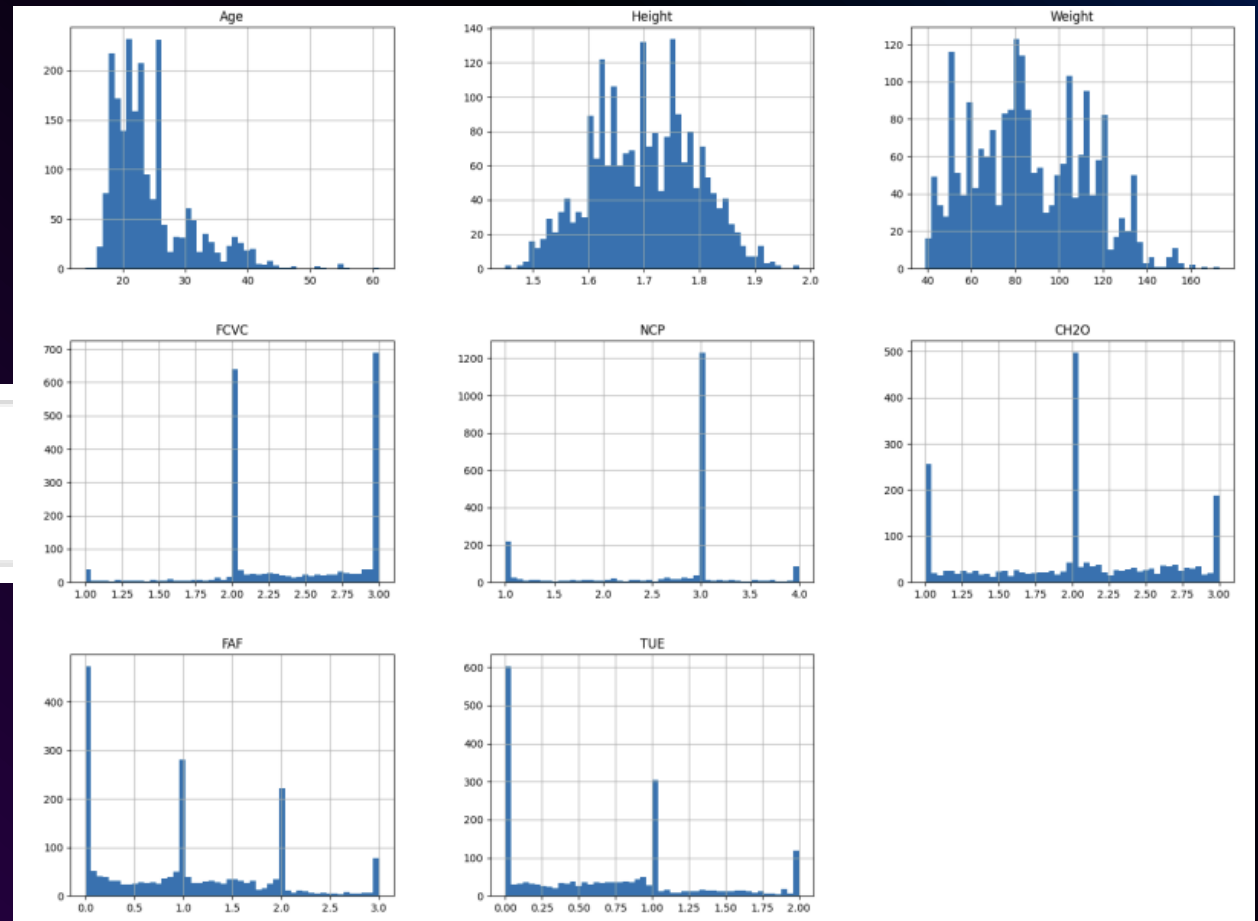
  

	CAEC	MTRANS	NOBeyesdad
0	Sometimes	Public_Transportation	Normal_Weight
1	Sometimes	Public_Transportation	Normal_Weight
2	Sometimes	Public_Transportation	Normal_Weight
3	Sometimes	Walking	Overweight_Level_I
4	Sometimes	Public_Transportation	Overweight_Level_II
...	...	...	...
2106	Sometimes	Public_Transportation	Obesity_Type_III
2107	Sometimes	Public_Transportation	Obesity_Type_III
2108	Sometimes	Public_Transportation	Obesity_Type_III
2109	Sometimes	Public_Transportation	Obesity_Type_III
2110	Sometimes	Public_Transportation	Obesity_Type_III

[2087 rows x 17 columns]

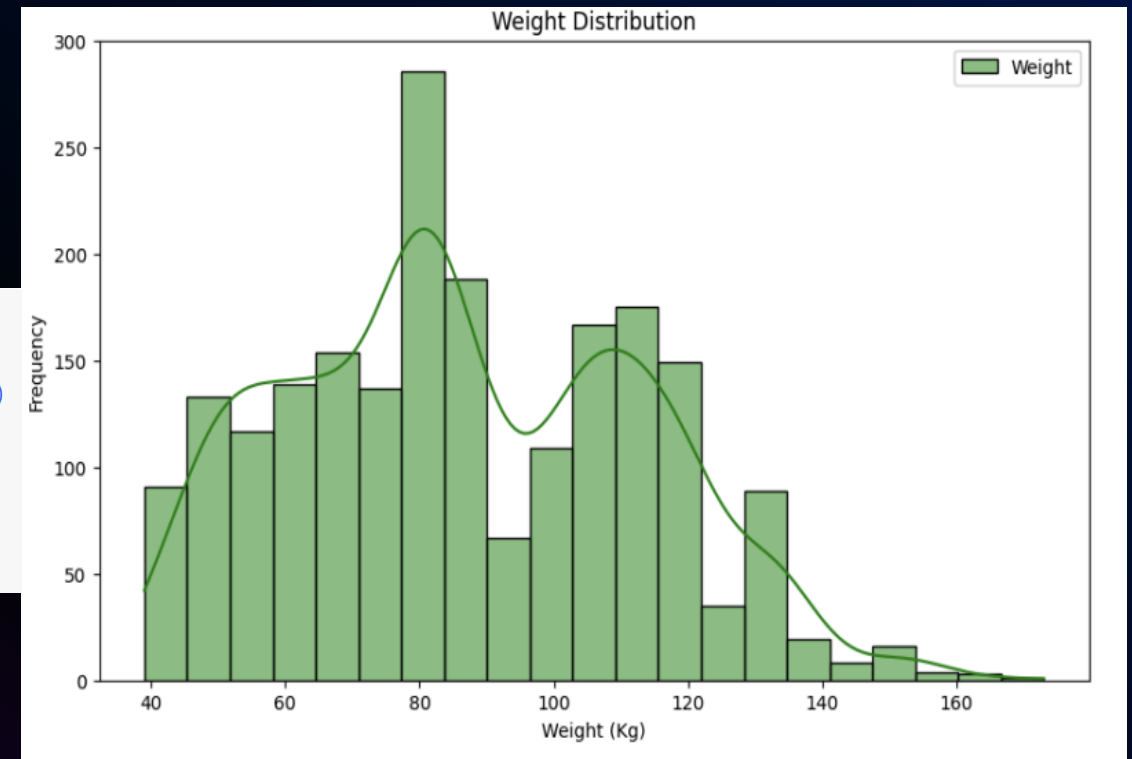
# GENERATING HISTOGRAM PLOTS FOR THE DATASET ATTRIBUTES

```
# Plot histograms for different parameters  
data.hist(bins=50, figsize=(20,15));
```



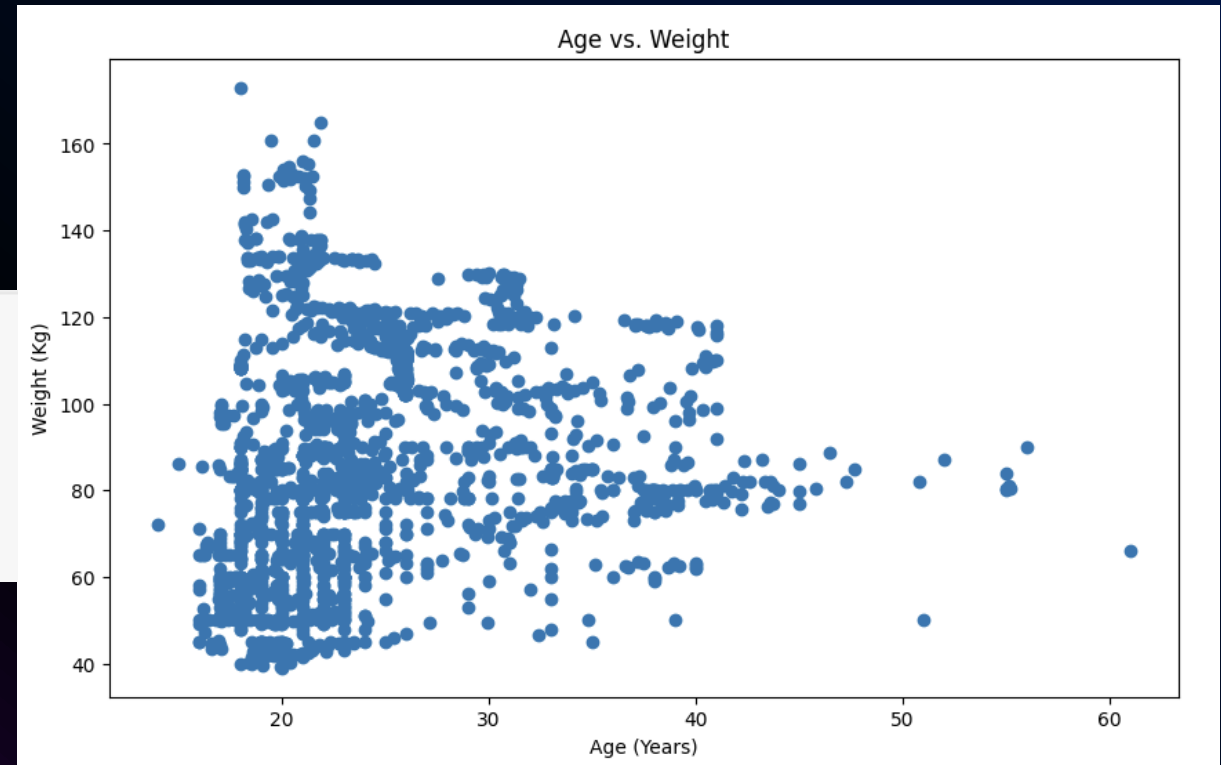
# HISTOGRAM PLOT SHOWING THE WEIGHT DISTRIBUTION

```
1 # Plot Weight Distribution
2 plt.figure(figsize=(10, 6))
3 sns.histplot(data=data, x='Weight', color='green', kde=True, label='Weight')
4 plt.title('Weight Distribution')
5 plt.xlabel('Weight (Kg)')
6 plt.ylabel('Frequency')
7 plt.legend()
8 plt.show()
```



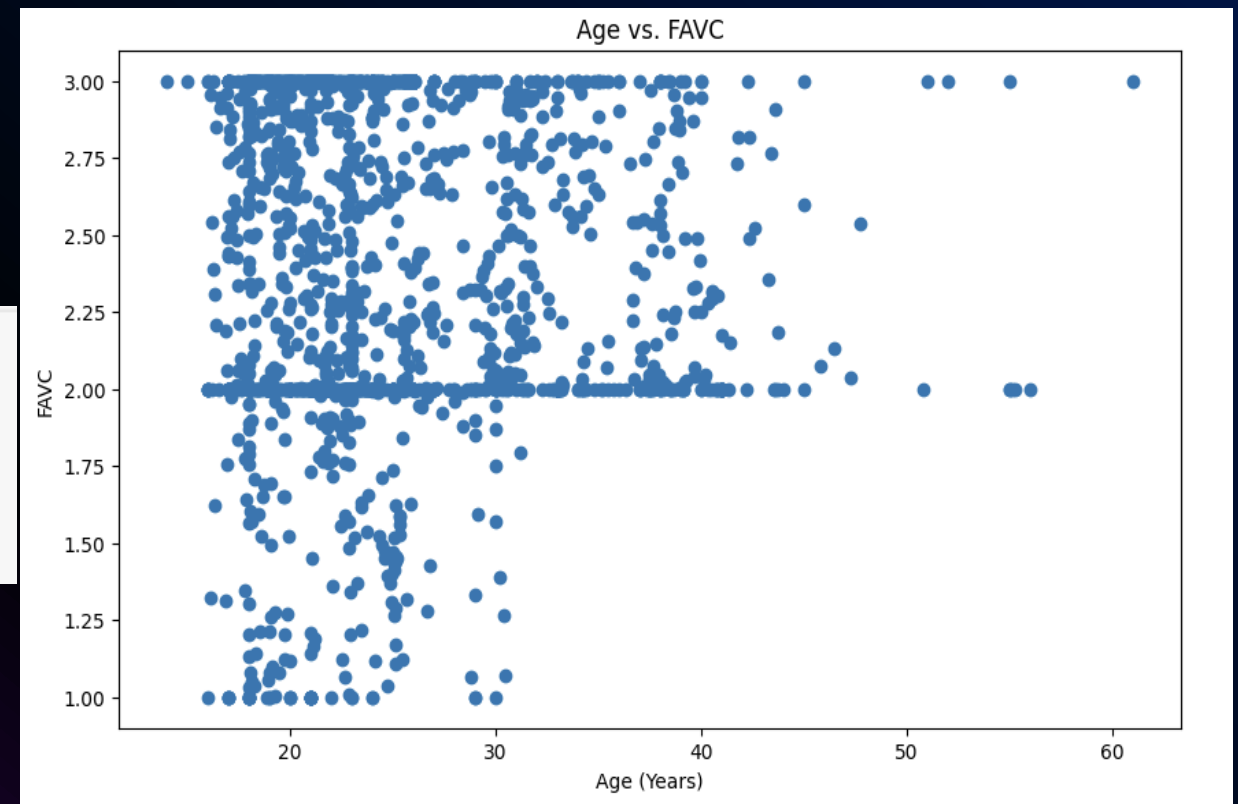
# SCATTER PLOT DEPICTING AGE VS. WEIGHT

```
1 fig, ax = plt.subplots(figsize=(10,6))
2 ax.scatter(data['Age'], data['Weight'])
3 plt.title('Age vs. Weight')
4 ax.set_xlabel('Age (Years)')
5 ax.set_ylabel('Weight (Kg)')
6 plt.show()
```



# SCATTER PLOT SHOWING AGE VS. FCVC

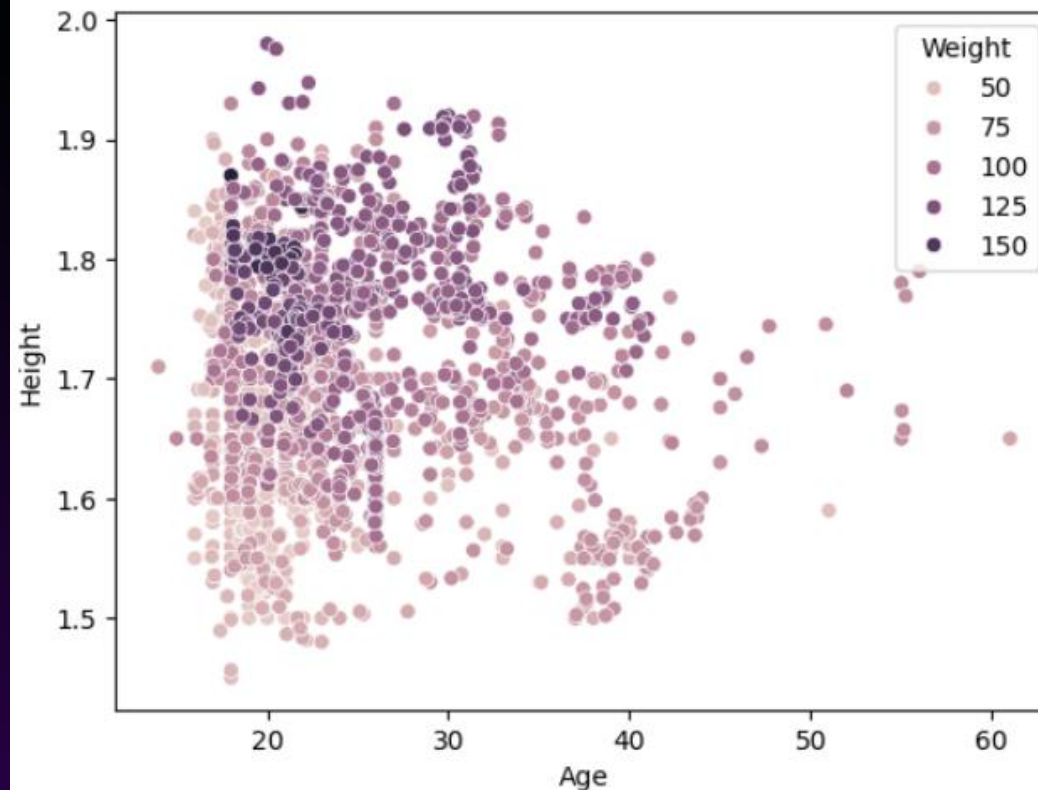
```
1 fig, ax = plt.subplots(figsize=(10,6))
2 ax.scatter(data['Age'], data['FCVC'])
3 plt.title('Age vs. FAVC')
4 ax.set_xlabel('Age (Years)')
5 ax.set_ylabel('FAVC')
6 plt.show()
```





# SCATTER PLOT ILLUSTRATING AGE VS. HEIGHT FOR DIFFERENT WEIGHTS

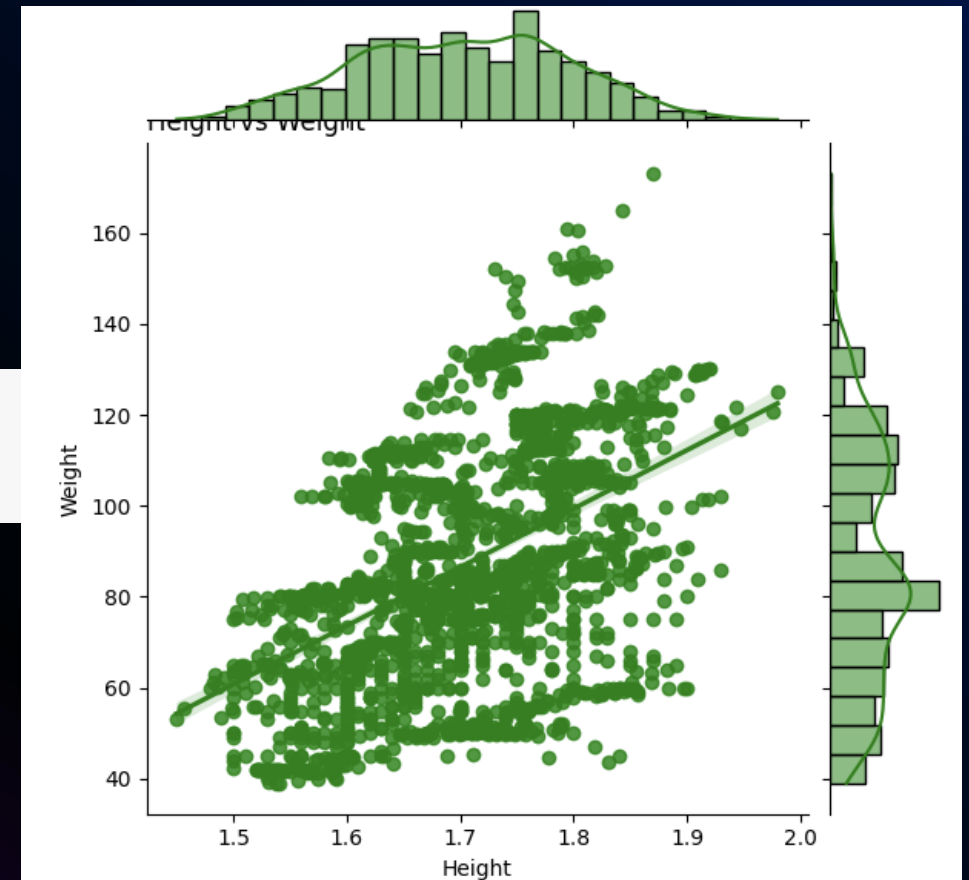
```
1 scatter_P = sns.scatterplot(x= "Age" ,y= "Height",
2                               hue="Weight",
3                               data=data);
```



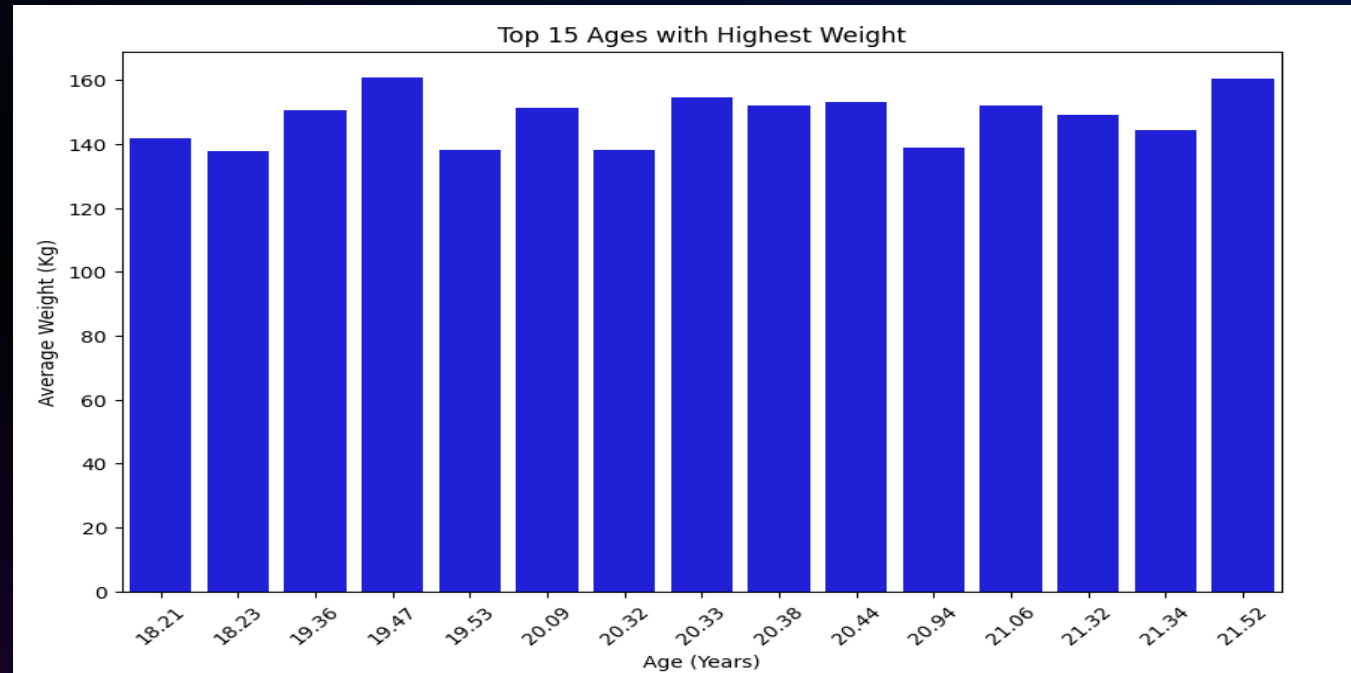


# USING 'JOINTPLOT' FOR PLOTTING HEIGHT VS. WEIGHT

```
1 sns.jointplot(x="Height",y='Weight',data=data,kind='reg', color='green')  
2 plt.title("Height vs Weight",loc='left')
```



# BAR PLOT SHOWING AGE VS. AVERAGE WEIGHT FOR THE TOP 15 AGES

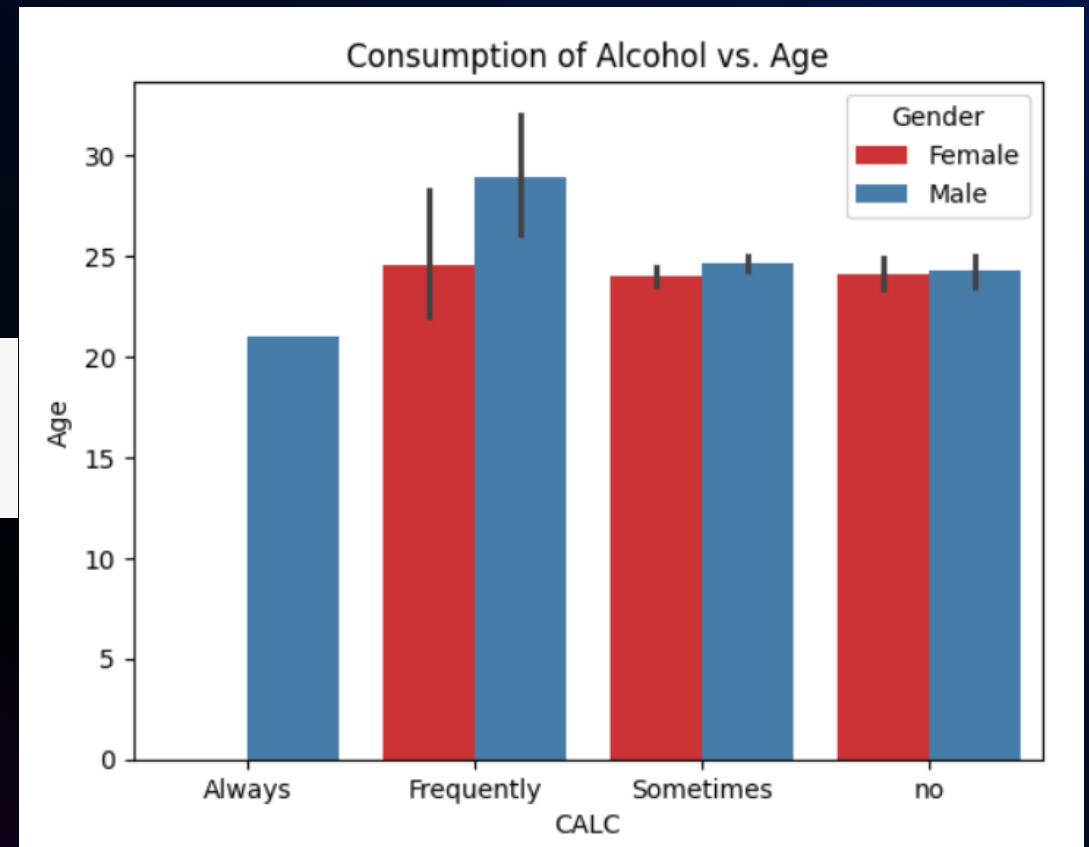


```
1 # Top 15 Ages with Highest Weight
2 top_15_ages = data.groupby('Age')['Weight'].mean().nlargest(15)
3 plt.figure(figsize=(10, 6))
4 sns.barplot(x=top_15_ages.index, y=top_15_ages.values, color='blue')
5 plt.title('Top 15 Ages with Highest Weight')
6 plt.xlabel('Age (Years)')
7 plt.ylabel('Average Weight (Kg)')
8 plt.xticks(rotation=45)
9 plt.show()
```

```
1 data['Age']=data['Age'].round(2)
```

# BARPLOT ILLUSTRATING CONSUMPTION OF ALCOHOL FOR FEMALES & MALES

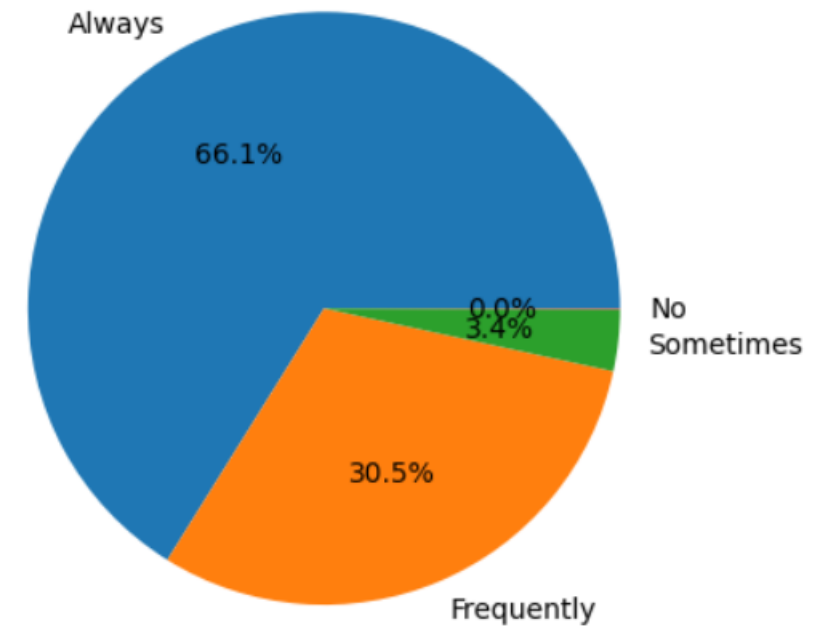
```
1 data['CALC'] = data['CALC'].astype('category')
2 sns.barplot(x='CALC', y='Age', data=data, hue='Gender', palette='Set1')
3 plt.title('Consumption of Alcohol vs. Age')
4 plt.show()
```



# PIE CHART SHOWING THE % CONSUMPTION OF ALCOHOL

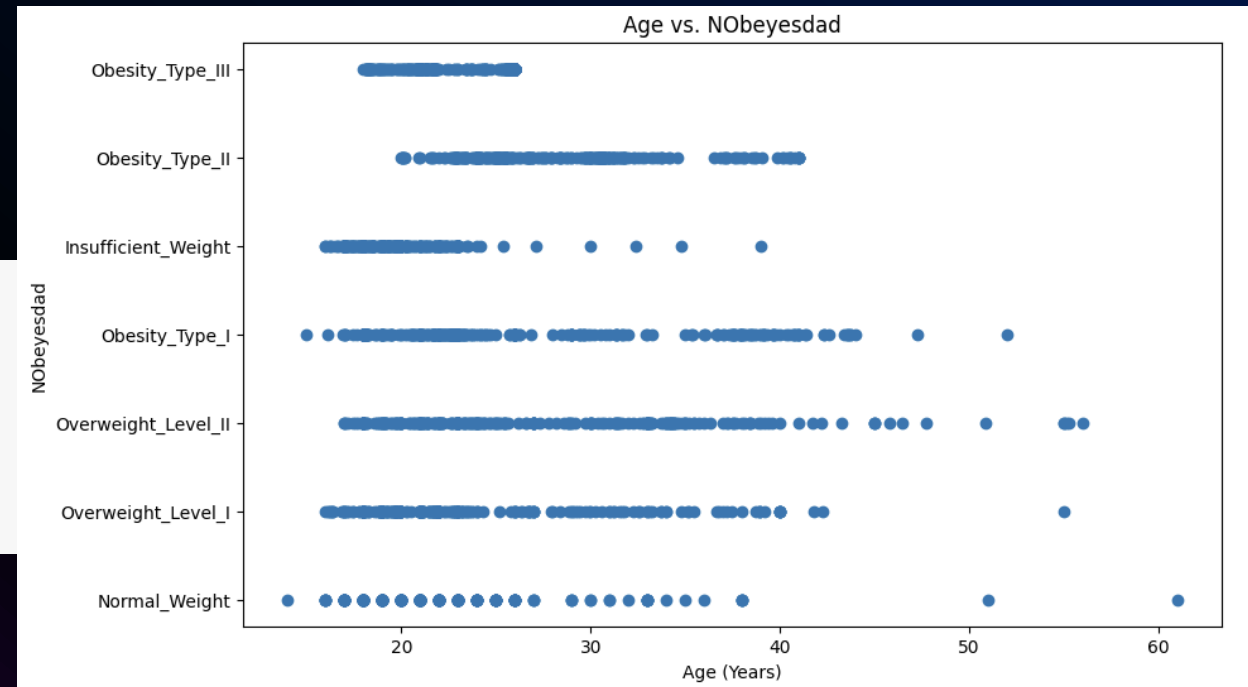
```
1 count = data['CALC'].value_counts()
2 labels = ["Always", "Frequently", "Sometimes", "No"]
3 vals = count.values
4 plt.pie(vals, labels=labels, autopct="%1.1f%%")
5 plt.title("% Consumption of Alcohol for Males & Females")
6
7 plt.show()
```

% Consumption of Alcohol for Males & Females



# PLOTTING AGE VS. NOBEYESDAD USING SCATTER PLOT

```
1 # Age vs. NObeyesdad
2 fig, ax = plt.subplots(figsize=(10,6))
3 ax.scatter(data['Age'], data['NObeyesdad'])
4 plt.title('Age vs. NObeyesdad')
5 ax.set_xlabel('Age (Years)')
6 ax.set_ylabel('NObeyesdad')
7 plt.show()
```

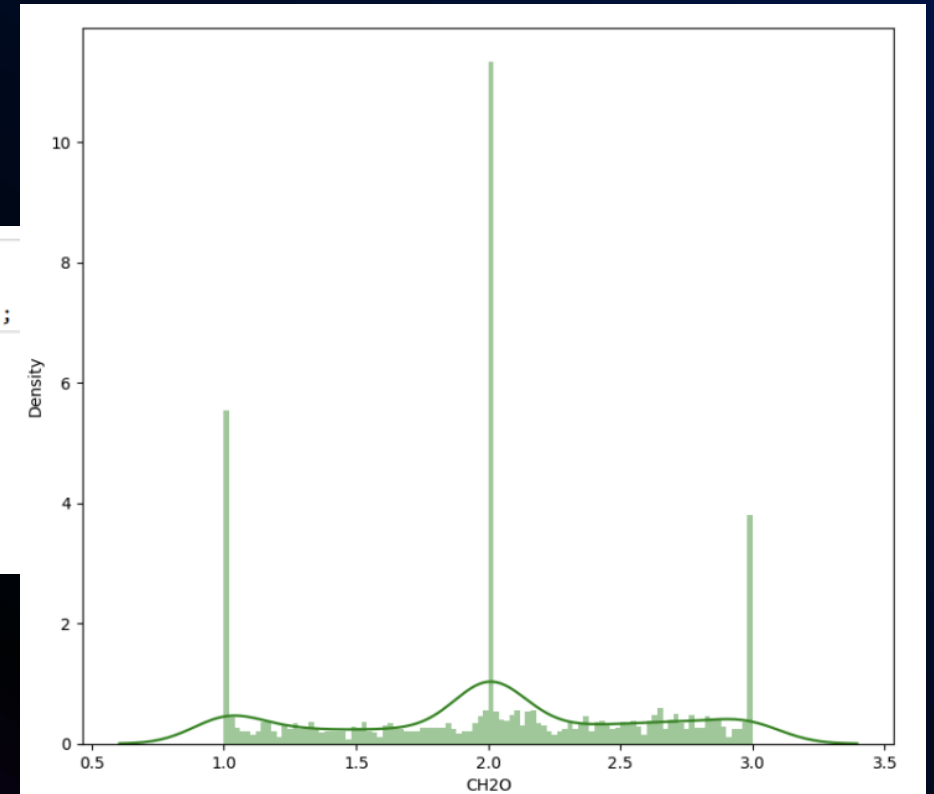


# PLOTTING THE DISTRIBUTION OF CH2O

```
print(data['CH2O'].describe())
plt.figure(figsize=(9, 8))
sns.distplot(data['CH2O'], color='g', bins=100, hist_kws={'alpha': 0.4});
```

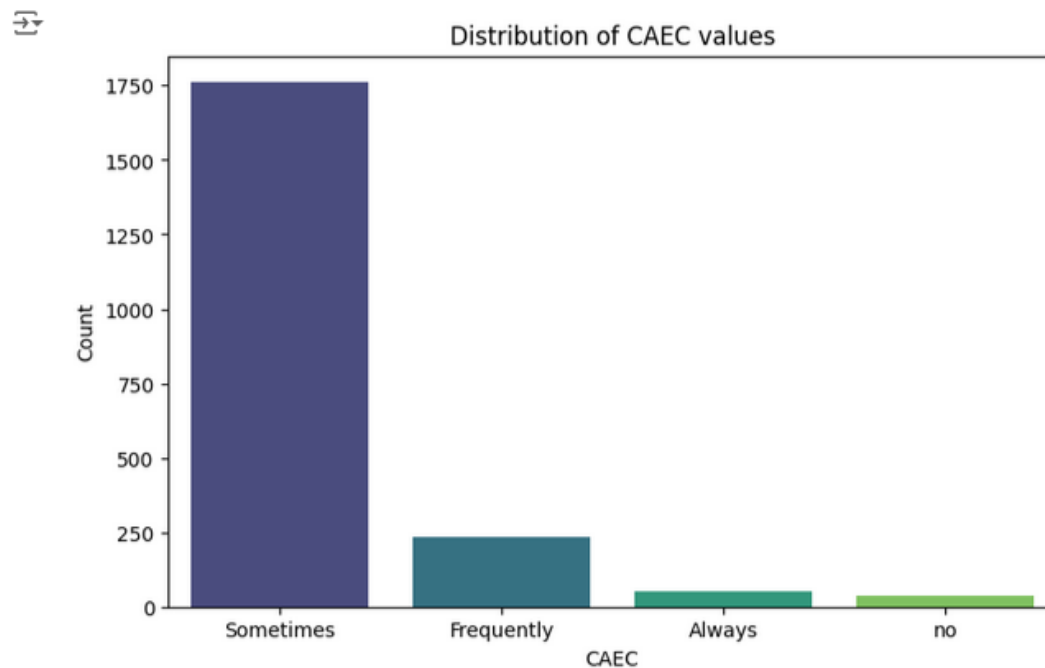
count	2087.000000
mean	2.004749
std	0.608284
min	1.000000
25%	1.590922
50%	2.000000
75%	2.466193
max	3.000000

Name: CH2O, dtype: float64



# COUNT PLOT SHOWING THE CATEGORICAL FACTOR 'CAEC'

```
1 # Distribution of CAEC values
2 plt.figure(figsize=(8, 5))
3 sns.countplot(data=data, x='CAEC', palette='viridis')
4 plt.title('Distribution of CAEC values')
5 plt.xlabel('CAEC')
6 plt.ylabel('Count')
7 plt.show()
8
```



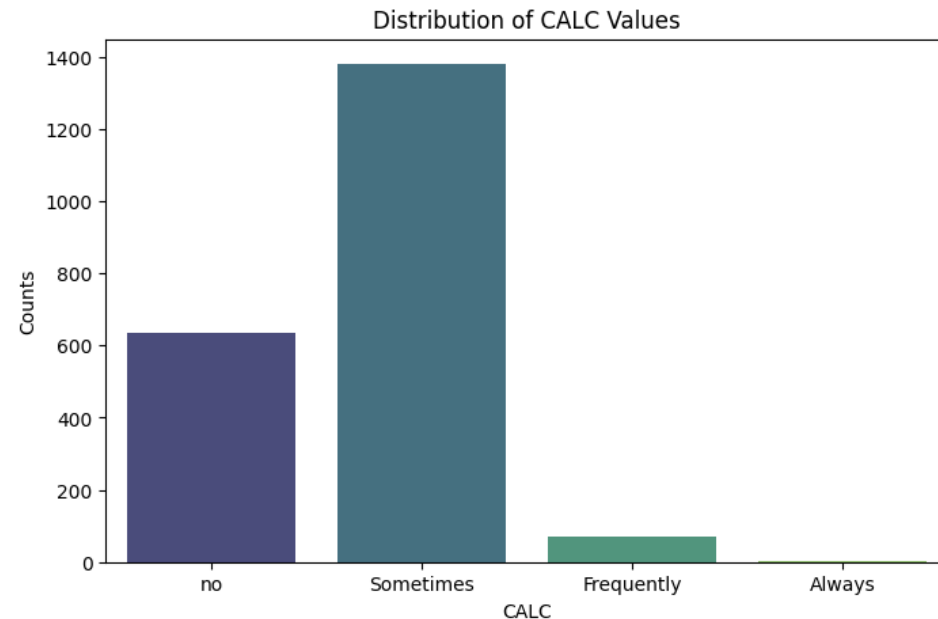


# COUNT PLOT DEPICTING THE CATEGORICAL FACTOR 'CALC'

```
1 data['CALC'].unique()
```

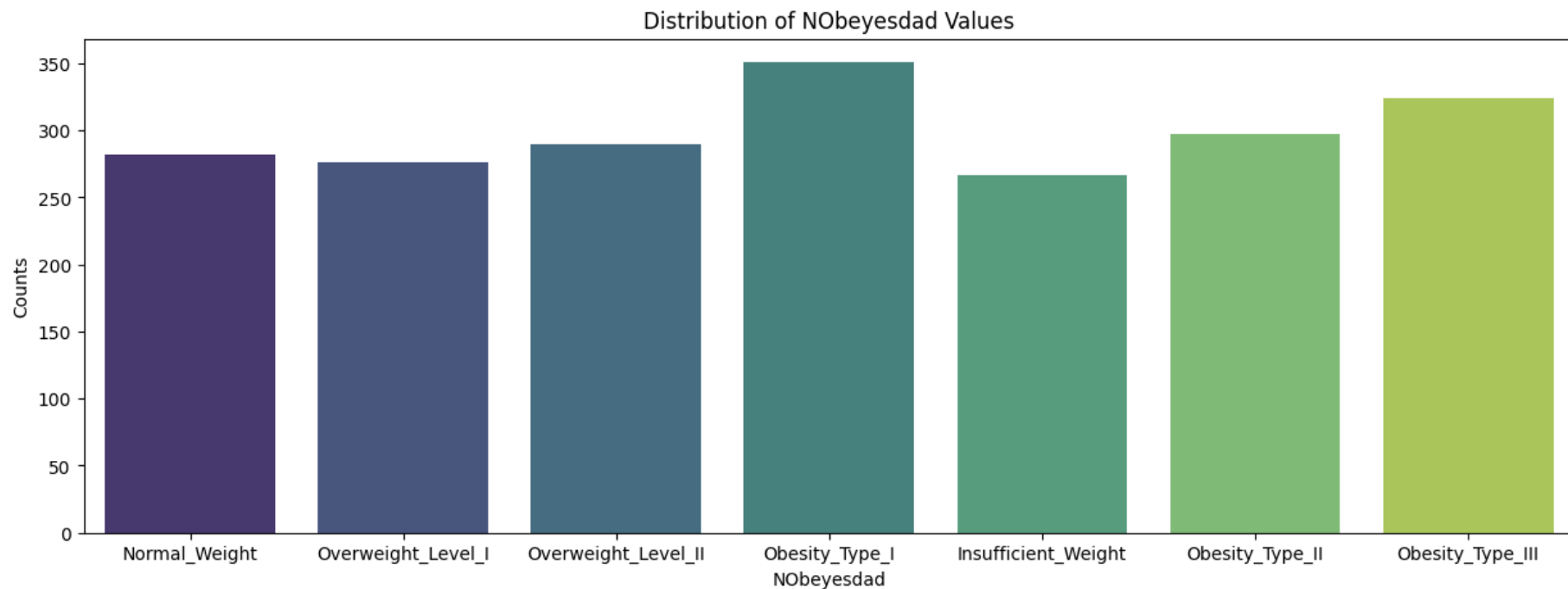
```
array(['no', 'Sometimes', 'Frequently', 'Always'], dtype=object)
```

```
1 # CALC distribution
2 plt.figure(figsize=(8,5))
3 sns.countplot(data=data, x='CALC', palette='viridis')
4 plt.title('Distribution of CALC Values')
5 plt.xlabel ('CALC')
6 plt.ylabel('Counts')
7 plt.show()
```



# COUNT PLOT DEPICTING NOBYESDAD VS. FREQUENCY

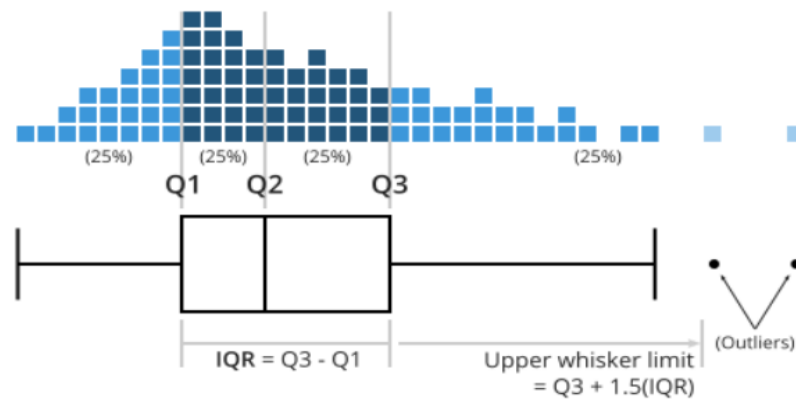
```
1 # NObeyesdad distribution
2 plt.figure(figsize=(15,5))
3 sns.countplot(data=data, x='NObeyesdad', palette='viridis')
4 plt.title('Distribution of NObeyesdad Values')
5 plt.xlabel ('NObeyesdad')
6 plt.ylabel('Counts')
7 plt.show()
```



# HOW TO INTERPRETING A BOX PLOT?

## Interpreting a box and whiskers

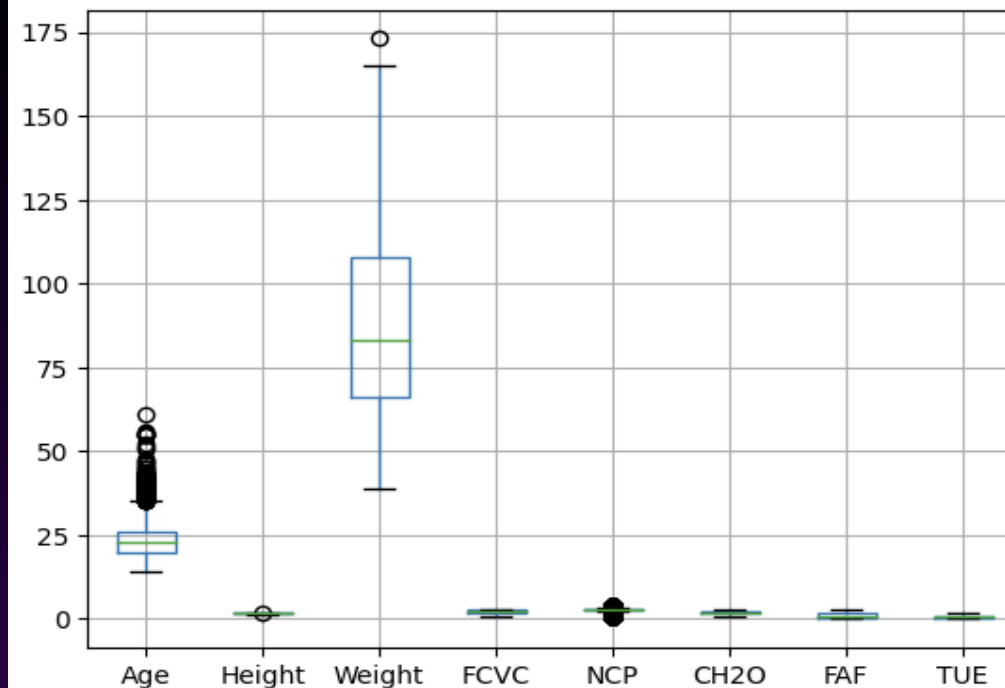
Construction of a box plot is based around a dataset's [quartiles](#), or the values that divide the dataset into equal fourths. The first quartile (Q1) is greater than 25% of the data and less than the other 75%. The second quartile (Q2) sits in the middle, dividing the data in half. Q2 is also known as the median. The third quartile (Q3) is larger than 75% of the data, and smaller than the remaining 25%. In a box and whiskers plot, the ends of the box and its center line mark the locations of these three quartiles.



<https://www.atlassian.com/data/charts/box-plot-complete-guide>

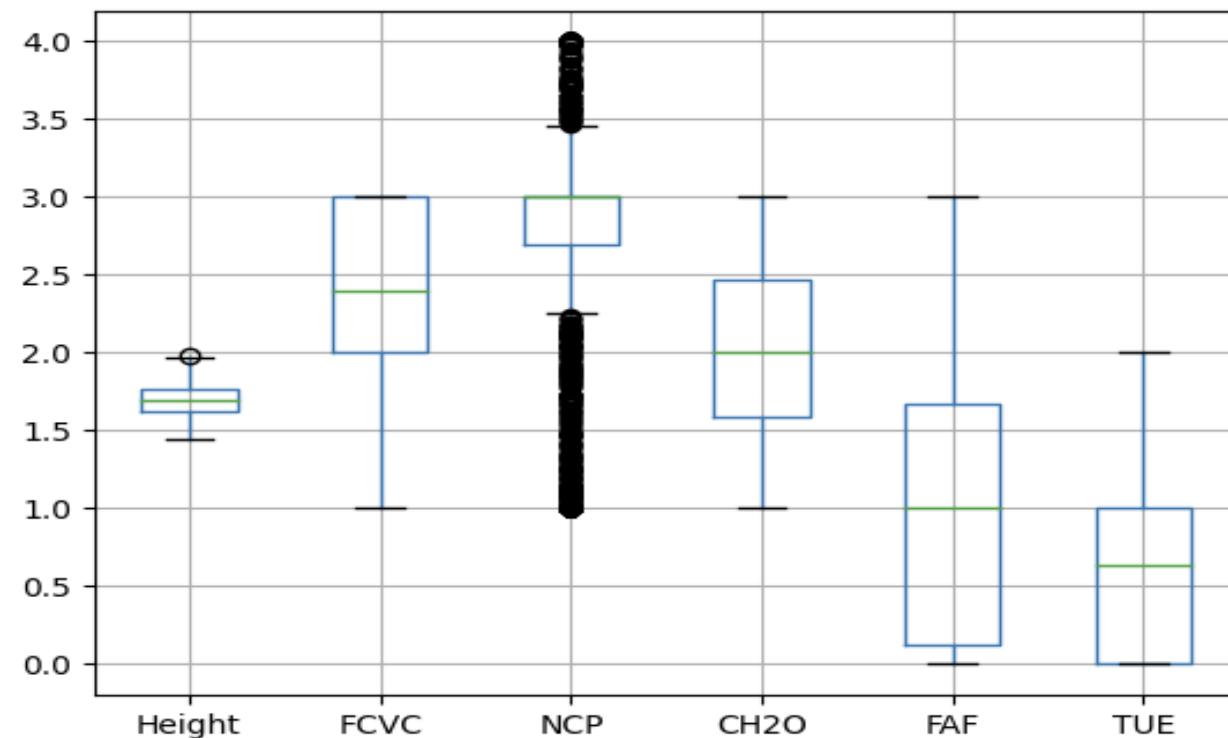
# INSPECTING THE DATASET FOR OUTLIERS USING BOXPLOT

```
1 columns=['Age', 'Height', 'Weight', 'FCVC', 'NCP', 'CH2O', 'FAF', 'TUE' ]  
2 boxplot = data.boxplot(column=['Age', 'Height', 'Weight', 'FCVC', 'NCP', 'CH2O', 'FAF', 'TUE'])
```

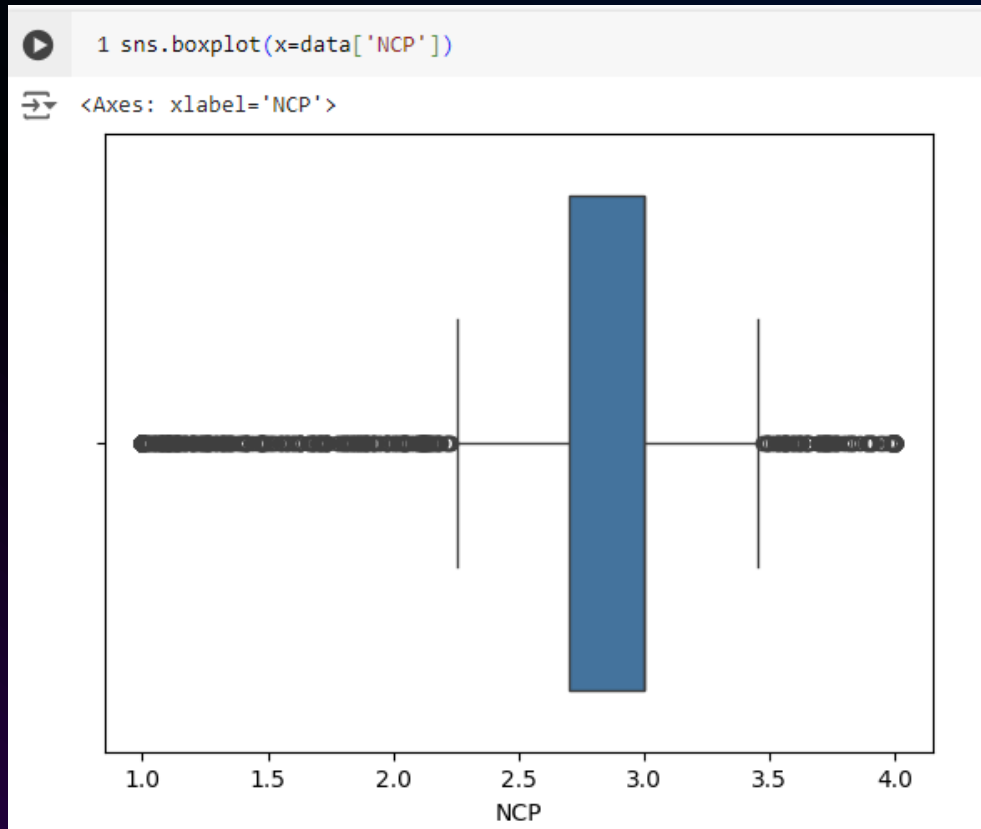


# INSPECTING THE DATASET FOR OUTLIERS USING BOXPLOT

```
1 columns=['Height', 'Weight', 'FCVC', 'NCP', 'CH2O', 'FAF', 'TUE' ]  
2 boxplot = data.boxplot(column=['Height', 'FCVC', 'NCP', 'CH2O', 'FAF', 'TUE'])
```



# BOXPLOT SHOWING THE NCP FACTOR VALUES



# THE DATA SET FACTORS BEFORE REMOVING THE OUTLIERS

```
[16] 1 # define path
      2 data_path = '/content/sample_data/diabetes.csv'
      3
      4 # import/load data into a newly created dataframe, df
      5 df = pd.read_csv(data_path)
```

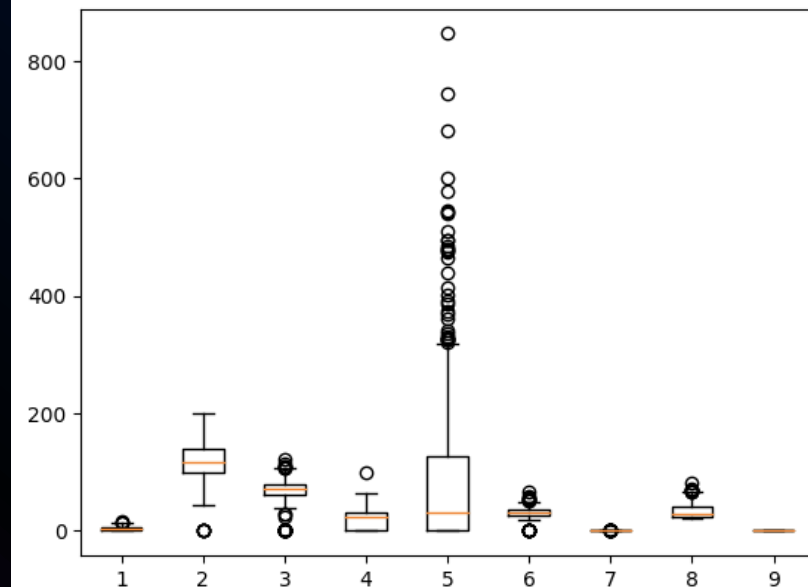
```
[17] 1 df.columns
```

```
Index(['Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness', 'Insulin',
       'BMI', 'DiabetesPedigreeFunction', 'Age', 'Outcome'],
      dtype='object')
```

## Data Set Attribute Information:

- Pregnancies
- Glucose
- BloodPressure
- SkinThickness
- Insulin
- BMI
- Diabetes Pedigree Function
- Age
- Outcome

```
1 plt.boxplot(df);
```



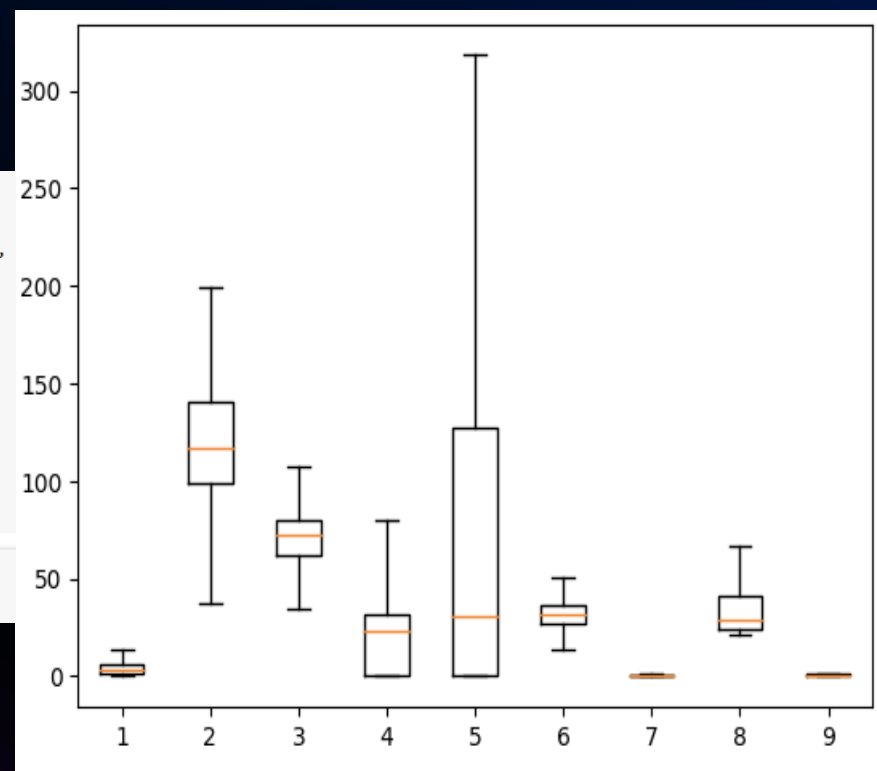
Example showing the removal of outliers using the diabetes.csv dataset  
<https://www.kaggle.com/datasets/saurabh00007/diabetescsv>



# THE DATASET FACTORS AFTER REMOVING THE OUTLIERS

```
[37] 1 # Treat outliers in the data set
      2 def outlier_removal():
      3     l = ['Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness', 'Insulin',
      4         'BMI', 'DiabetesPedigreeFunction', 'Age', 'Outcome']
      5     for i in l:
      6         x = np.quantile(df[i], [0.25, 0.75])
      7         iqr = x[1] - x[0]
      8         upper = x[1] + 1.5 * iqr
      9         lower = x[0] - 1.5 * iqr
     10         df[i] = np.where(df[i] > upper, upper, (np.where(df[i] < lower, lower, df[i])))
     11
     12 outlier_removal()
     13
```

```
1 # Plot data after removing outliers
2 plt.boxplot(df);
```



<https://www.geeksforgeeks.org/detect-and-remove-the-outliers-using-python/>

# WHAT IS A CORRELATION?

Correlation is a statistical indicator that quantifies the degree to which two variables change in relation to each other. It indicates the strength and direction of the linear relationship between two variables. The correlation coefficient is denoted by “r”, and it ranges from -1 to 1.

- If  $r = -1$ , it means that there is a perfect negative correlation.
- If  $r = 0$ , it means that there is no correlation between the two variables.
- If  $r = 1$ , it means that there is a perfect positive correlation.

There are two popular methods used to find the correlation coefficients:

## **Pearson's product-moment correlation coefficient**

The Pearson correlation coefficient ( $r$ ) is a measure of linear relationship between two variables.

$$r = \frac{n(\sum xy) - (\sum x)(\sum y)}{\sqrt{[n \sum x^2 - (\sum x)^2][n \sum y^2 - (\sum y)^2]}}$$

Here,

- $n$  is the number of data points
- $\sum xy$  is the sum of the product of corresponding values of  $x$  and  $y$
- $\sum x$  is the sum of all the values of  $x$
- $\sum y$  is the sum of all the values of  $y$
- $\sum x^2$  is the sum of the squares of all values of  $x$
- $\sum y^2$  is the sum of the squares of all the of  $y$

# WHAT IS A CORRELATION MATRIX?

---

A correlation is a tabular representation that displays correlation coefficients, indicating the strength and direction of relationships between variables in a dataset. Within this matrix, each cell signifies the correlation between two specific variables. This tool serves multiple purposes, serving as a summary of data relationships, input for more sophisticated analyses, and a diagnostic aid for advanced analytical procedures. By presenting a comprehensive overview of inter-variable correlations, the matrix becomes invaluable in discerning patterns, guiding further analyses, and identifying potential areas of interest or concern in the dataset. Its applications extend beyond mere summary statistics, positioning it as a fundamental component in the preliminary stages of diverse and intricate data analyses.

<https://www.geeksforgeeks.org/create-a-correlation-matrix-using-python/>

# INTERPRETING THE CORRELATION MATRIX RESULTS

---

Strong correlations, indicated by values close to 1 or -1, suggest a robust connection, while weak correlations, near 0, imply a less pronounced association. They are identifying these degrees of correlation aids in understanding the intensity of interactions within the dataset, facilitating targeted analysis and decision-making. Positive correlations (values  $> 0$ ) signify that as one variable increases, the other tends to increase as well. Conversely, negative correlations (values  $< 0$ ) imply an inverse relationship—when one variable increases, the other tends to decrease. Investigating these directional associations provides insights into how variables influence each other, crucial for formulating informed hypotheses and predictions.

<https://www.geeksforgeeks.org/create-a-correlation-matrix-using-python/>

# CALCULATING THE PAIRWISE CORRELATION FOR ALL COLUMNS

```
[ ] 1 numeric_df = data.select_dtypes(include=['number'])# Select only numeric columns
    2 # Calculate correlation matrix
    3 correlation_matrix = numeric_df.corr()
    4 # Print the correlation matrix
    5 print("Correlation Matrix:")
    6 print(correlation_matrix)
```

Correlation Matrix:

	Age	Height	Weight	FCVC	NCP	CH2O	FAF	\
Age	1.000000	-0.031748	0.198160	0.013572	-0.055823	-0.044058	-0.148202	
Height	-0.031748	1.000000	0.457468	-0.040363	0.227806	0.220487	0.293584	
Weight	0.198160	0.457468	1.000000	0.216574	0.092149	0.203823	-0.056490	
FCVC	0.013572	-0.040363	0.216574	1.000000	0.034885	0.081332	0.022003	
NCP	-0.055823	0.227806	0.092149	0.034885	1.000000	0.075335	0.127816	
CH2O	-0.044058	0.220487	0.203823	0.081332	0.075335	1.000000	0.165310	
FAF	-0.148202	0.293584	-0.056490	0.022003	0.127816	0.165310	1.000000	
TUE	-0.302927	0.041808	-0.079351	-0.104128	0.015693	0.020704	0.058716	

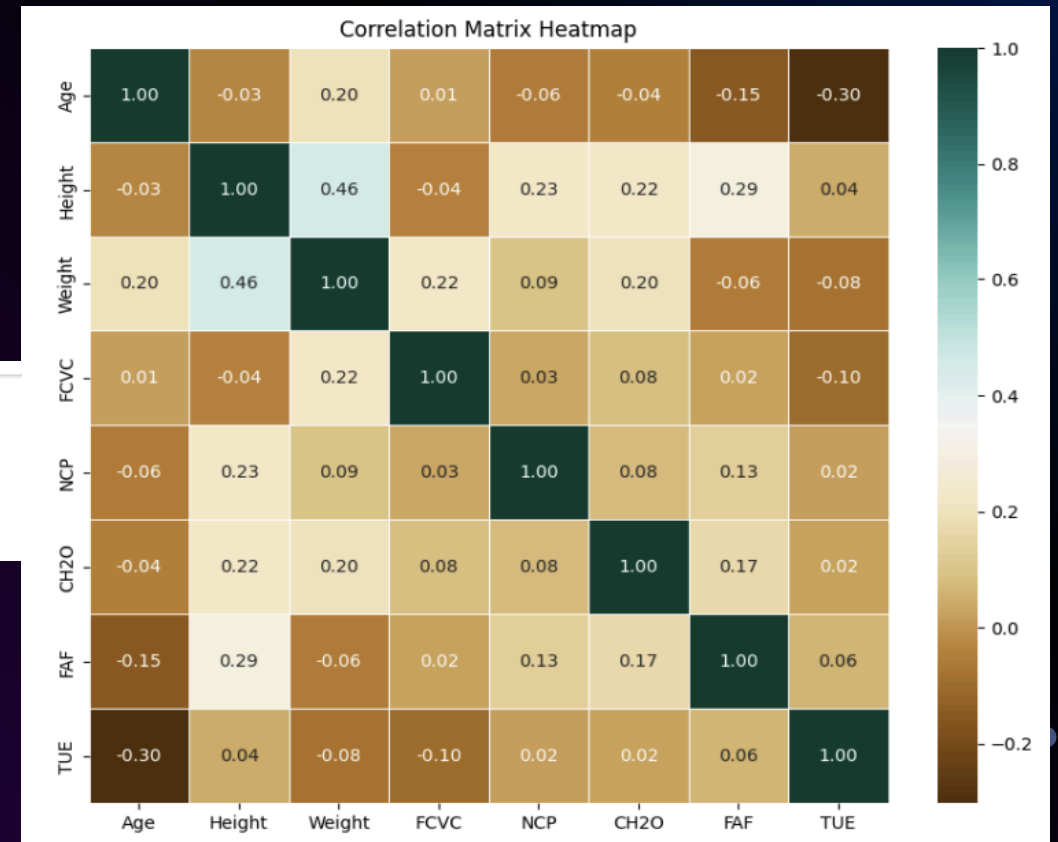
  

	TUE
Age	-0.302927
Height	0.041808
Weight	-0.079351
FCVC	-0.104128
NCP	0.015693
CH2O	0.020704
FAF	0.058716
TUE	1.000000

<https://www.geeksforgeeks.org/python-pandas-dataframe-corr/>

# PLOTTING THE CORRELATION MATRIX HEATMAP

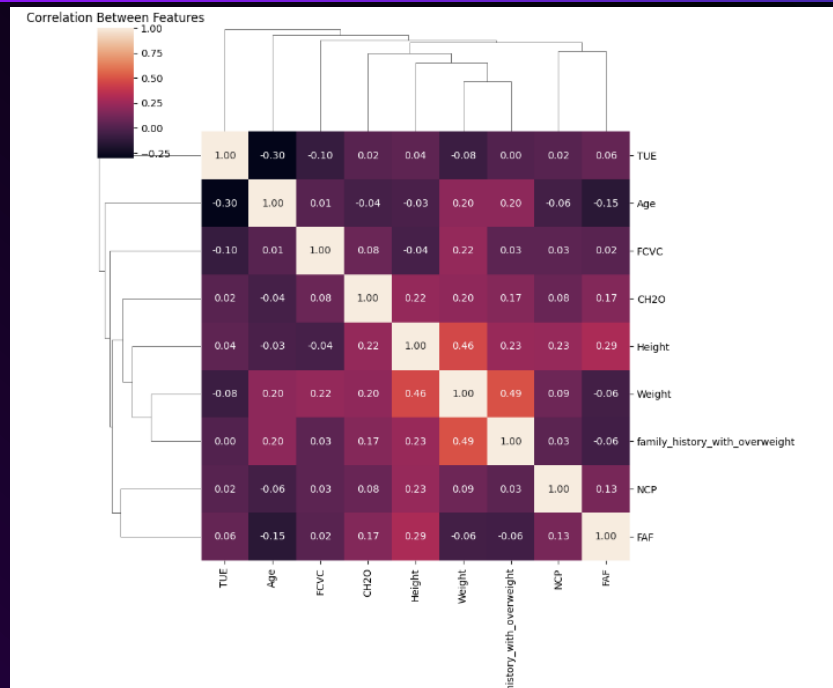
```
# Print the heatmap
plt.figure(figsize=(10, 8))
sns.heatmap(correlation_matrix, annot=True, cmap='BrBG', fmt=".2f", linewidths=0.5)
plt.title('Correlation Matrix Heatmap')
plt.show()
```



<https://www.geeksforgeeks.org/create-a-correlation-matrix-using-python/>



# CORRELATION MATRIX CLUSTER MAP



```
1 numeric_df = data.select_dtypes(include=['number'])# Select numeric columns only
2 # Calculate correlation matrix
3 correlation_matrix = numeric_df.corr()
4 sns.clustermap(correlation_matrix, annot=True, fmt=".2f")
5 plt.title("Correlation Between Features")
```



# After



1

# CONCLUSIONS

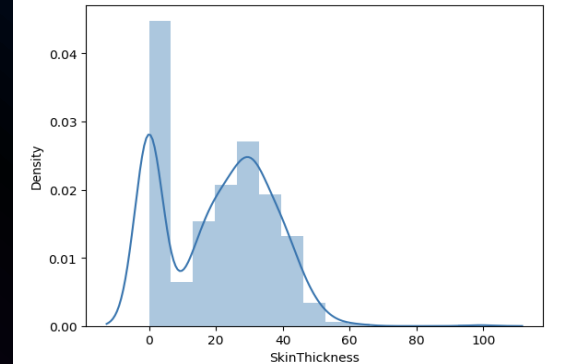
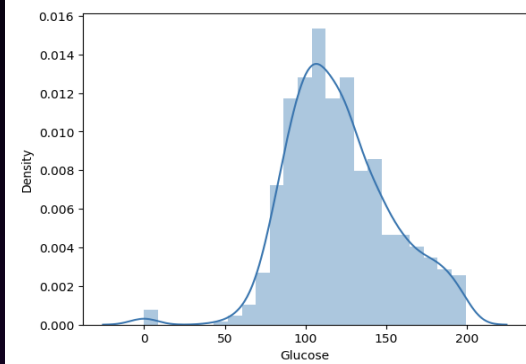
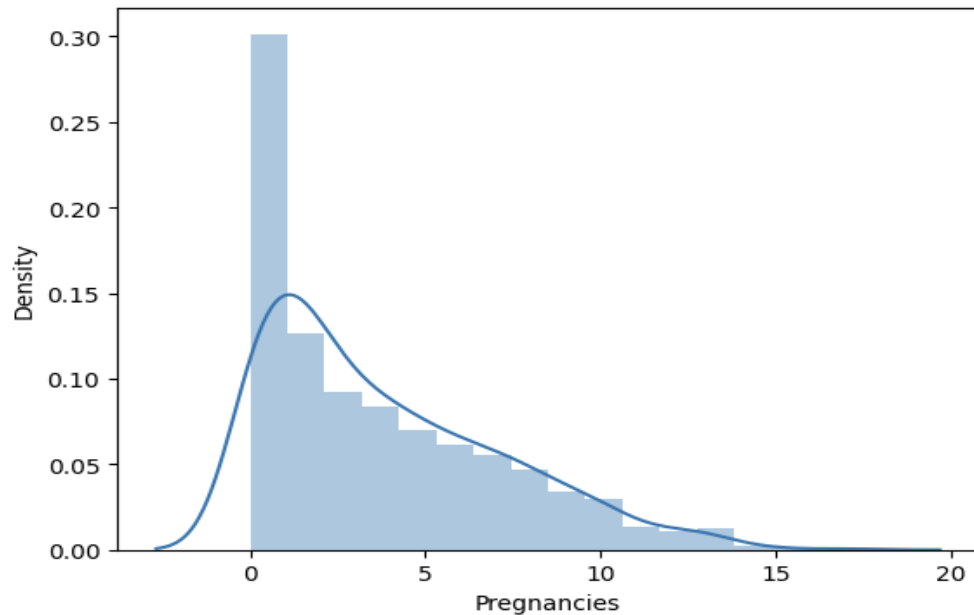
---

- **The original dataset contains 2111 rows and 17 column**
- **There are 24 duplicate rows in the dataset**
- **All values in the dataset are unique and contain no null, NaN,  $\pm \infty$  or missing values**
- **The NCP (number of main meals per day) data contains some outlier values and required removal**
- **There is a significant correlation between weight and height**

# ADDITIONAL EXAMPLES

# DISTRIBUTION PLOT ILLUSTRATION

```
1 data_feature = data.columns
2
3 for feature in data_feature:
4     p = sns.distplot(a = data[feature])
5     plt.show()
```

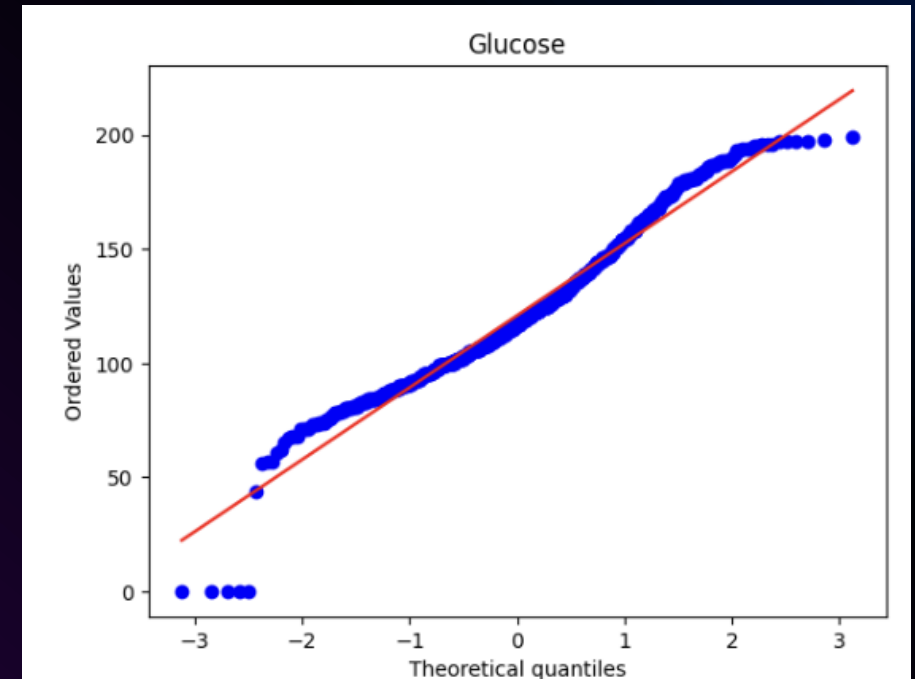
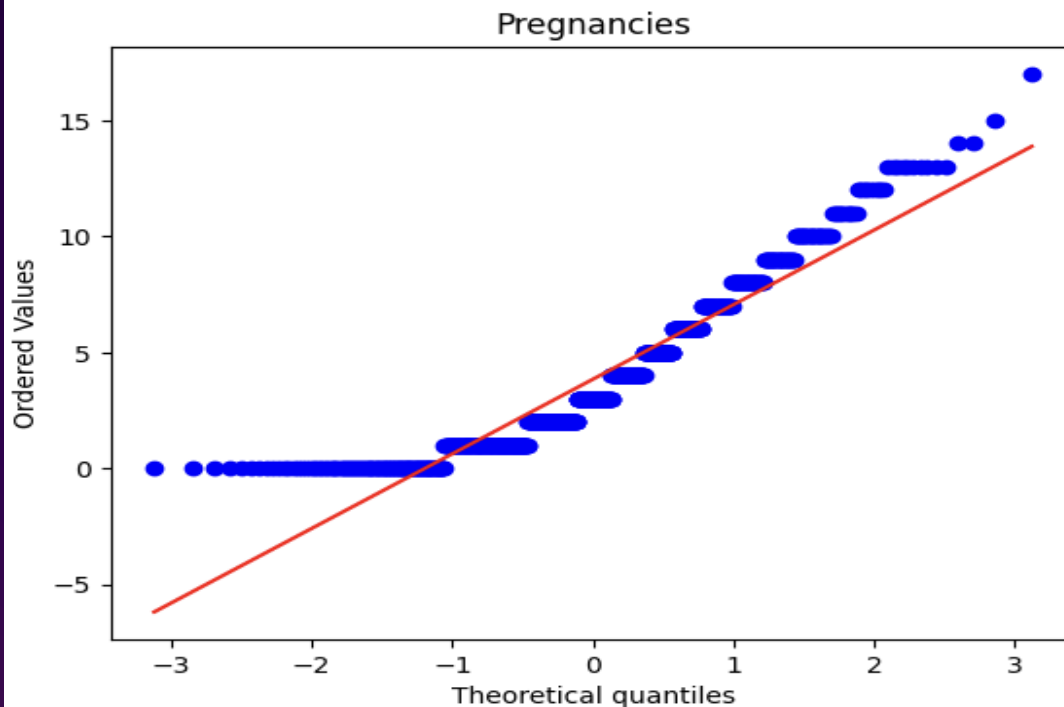


```
1 data = pd.read_csv("/content/sample_data/diabetes.csv") # Load data set using pandas
2 data.head(3)
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome
0	6	148	72	35	0	33.6	0.627	50	1
1	1	85	66	29	0	26.6	0.351	31	0
2	8	183	64	0	0	23.3	0.672	32	1

# PROBABILITY PLOT FOR TWO FACTORS IN THE DATASET

```
1 import scipy
2 from scipy import stats
3 for feature in data.columns:
4     stats.probplot(data[feature], plot = plt)
5     plt.title(feature)
6     plt.show()
```

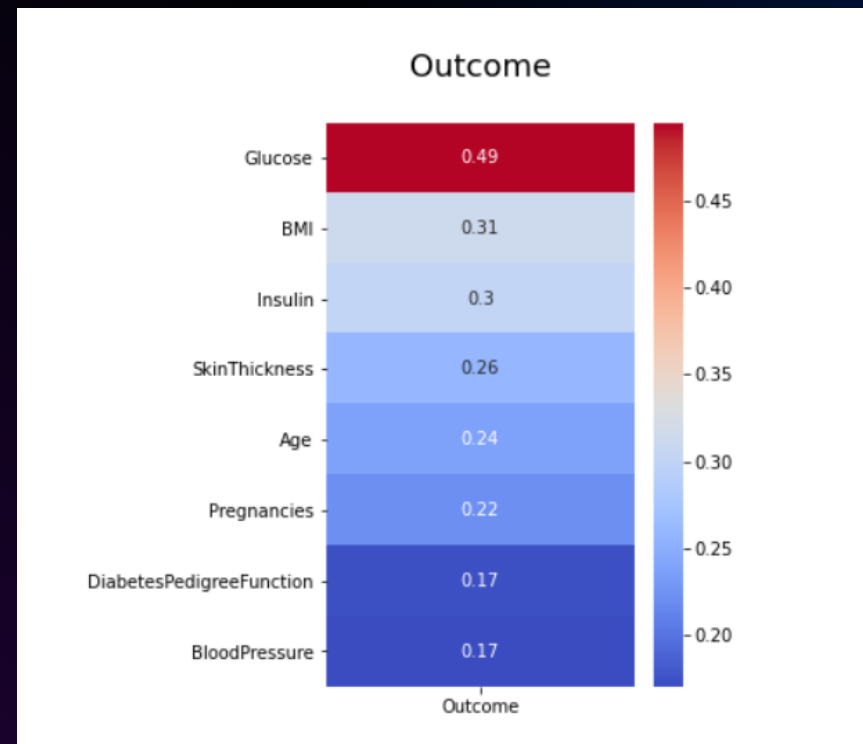


<https://www.kaggle.com/datasets/mathchi/diabetes-data-set>



# ANOTHER FORM OF HEATMAP PRESENTATION

```
def corr_to_target(dataframe, target, title=None, file=None):  
    plt.figure(figsize=(4,6))  
    sns.heatmap(dataframe.corr()[[target]].sort_values(target,  
                                                         ascending=False)  
[1:],  
               annot=True,  
               cmap='coolwarm')  
  
    plt.title(f'\n{title}\n', fontsize=18)  
  
    plt.show();  
  
    return  
  
corr_to_target(df, "Outcome", title="Outcome")
```



<https://www.kaggle.com/code/busekseolu/diabetes-classification>

# REFERENCES

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1. **Data Science Horizon, Data cleaning and preprocessing for data science beginners.**  
(<https://datasciencehorizons.com/data-cleaning-preprocessing-data-science-beginners-ebook/>)
2. **Matthieu Komorowski, et al., Exploratory Data Analysis, Chapter 15, doi:10.1007/978-3- 319-43742-2\_15.**
3. **DataQuest, Data Science Cheat Sheet-Pandas.** (<https://s3.amazonaws.com/dq-blog-files/pandas-cheat-sheet.pdf>)
4. **DataCamp, Python for Data Science Cheat Sheet-Matplotlib.**  
([https://s3.amazonaws.com/assets.datacamp.com/blog\\_assets/Python\\_Matplotlib\\_Cheat\\_Sheet.pdf](https://s3.amazonaws.com/assets.datacamp.com/blog_assets/Python_Matplotlib_Cheat_Sheet.pdf))
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([https://assets.datacamp.com/blog\\_assets/Numpy\\_Python\\_Cheat\\_Sheet.pdf](https://assets.datacamp.com/blog_assets/Numpy_Python_Cheat_Sheet.pdf))
6. **DataCamp, Python for Data Science, Seaborn Cheat Sheet.**  
([https://s3.amazonaws.com/assets.datacamp.com/blog\\_assets/Python\\_Seaborn\\_Cheat\\_Sheet.pdf](https://s3.amazonaws.com/assets.datacamp.com/blog_assets/Python_Seaborn_Cheat_Sheet.pdf))
7. **DataCamp, Python for Data Science Cheat Sheet-Scikit-Learn.**  
([https://s3.amazonaws.com/assets.datacamp.com/blog\\_assets/Scikit\\_Learn\\_Cheat\\_Sheet\\_Python.pdf](https://s3.amazonaws.com/assets.datacamp.com/blog_assets/Scikit_Learn_Cheat_Sheet_Python.pdf))
8. **David Beazley, et al., Python Cookbook, 3<sup>rd</sup> Ed., O'Reilly, Beijing, 2013.**